

Fig. 1

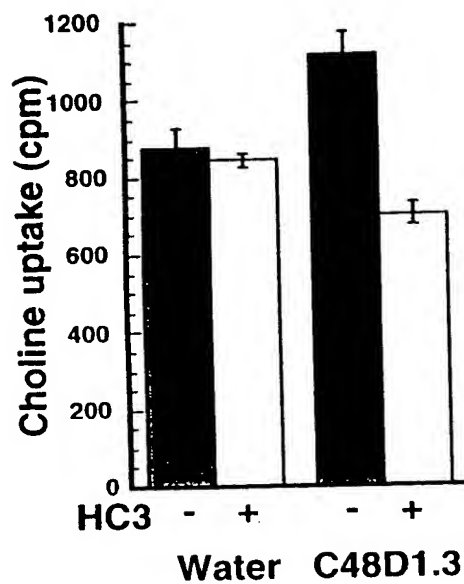


Fig. 2

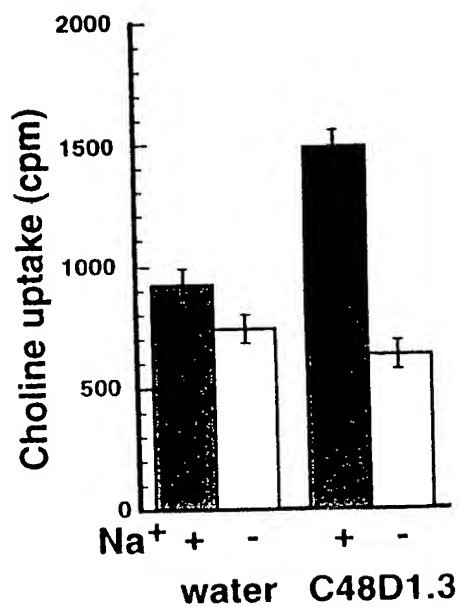


Fig. 3

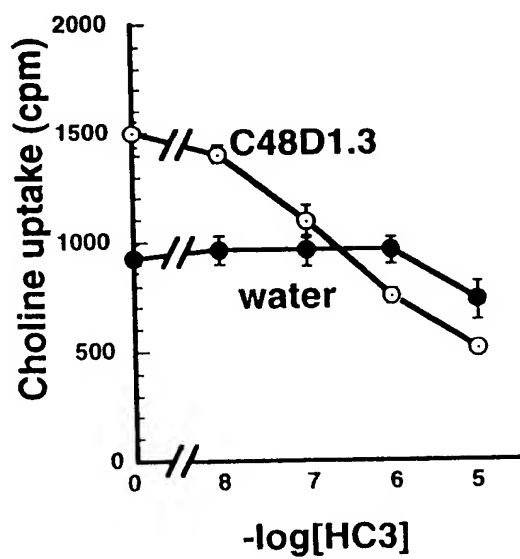


Fig. 4

CHT1	MPFHVEGVVAIDFYLLIVGIWAANKNS-----GNAEERSEAIVGGRIIGLLVGGF	56
cht-1	-MADLGVAIVFYLLIVGIWAGRKSSKELESAGAAIEEVLAGRNIIGLLVGIIF	59
<b>I</b>		
CHT1	TMTATWVGGGYINGTAEAYGPGCGLAWAQAPGYSLGGLFAKPMRSGYTMLD	116
cht-1	TMTATWVGGAINGTAEALYNGGLLGQAPGYSLGGLFAKPMREGYTMLD	117
<b>II</b>		
CHT1	PFQIKYGRYGGLLPALMGFWAAATISALGATISVIGDYNISVITSAITALLYT	176
cht-1	PFQIKYGRYGGLLPALMGFWTAATISALGATISVIGDYNASVITSAITALLYT	177
<b>III</b>		
CHT1	IVGGLYVAYTDVVQLFCIFGLWISVPFALSHPVVDIGFTAVHAKYQSPWLTIES-V	235
cht-1	ITGGYVAYTDVVQLFCIFGLWCVPAAVHDGAKDTSRNAG-----DWLGEITGGFK	231
<b>V</b>		
CHT1	EYVTWLDNALLLVGGIPWQAYFQVRLSSSATYAQVLSFAAFGCCMAIPAICIGATG	295
cht-1	ETSLWIDCILLLVGGIPWQVYFQVRLSSKAHGAQTLSEVAGVGCMAIPALIGATA	291
<b>VI</b>		
CHT1	ASTDWNQTAAYGFPDPKTKKEAD-----MILPEVLYQYLCPVYISFGLGAVSAAMSSAD	349
cht-1	RNTDWRMTDYSPWNGTKVESIPDPKRMVPEVLYQYLTPRIYFGLGAVSAAMSSAD	351
<b>VII</b>		
CHT1	SSLLSASSMFARNIYQLSTRQNASPKELVWMRIIVVFGASATAMALLTQVYGLWYLS	409
cht-1	SSLLSASSMFARNIYQLSTRQNASPKELVWMRIIVVFGIMATIMALTIDQVYGLWYLC	411
<b>VIII</b>		
CHT1	DLVYITFPQLLCVIRKGINTYGAVAGYIFGLFLRITGGEPYLYLPFYFGYYPDK	469
cht-1	DLVYITFPQLLCVIRPRNTYGSAGYAVGLVLRITGGEPYLYLPFYHYPMYT-D	469
<b>IX</b>		
CHT1	NGIYNQRFPPKTLTSMVISEFTNICVSYLAKYLFESGTLPPKLDITDAVVSRR---HSEENM	526
cht-1	G---VOYFPFRITAMSSMATIYIVSIQSEKLFSGRLSPWDYMGCVNIPIDHVPLPS	526
<b>X</b>		
CHT1	DKITLVRNENIKLNELAPVKPRQSTLSSTFTNKEALLDVDSSPEGSGTIDNLQ	580
cht-1	DVSTAVSSE---TLNMKAPNGTPAPVHPNQPSDENITLLHPYSDQSYYSITSN--	576
<b>XI</b>		
<b>XII</b>		

Fig. 5

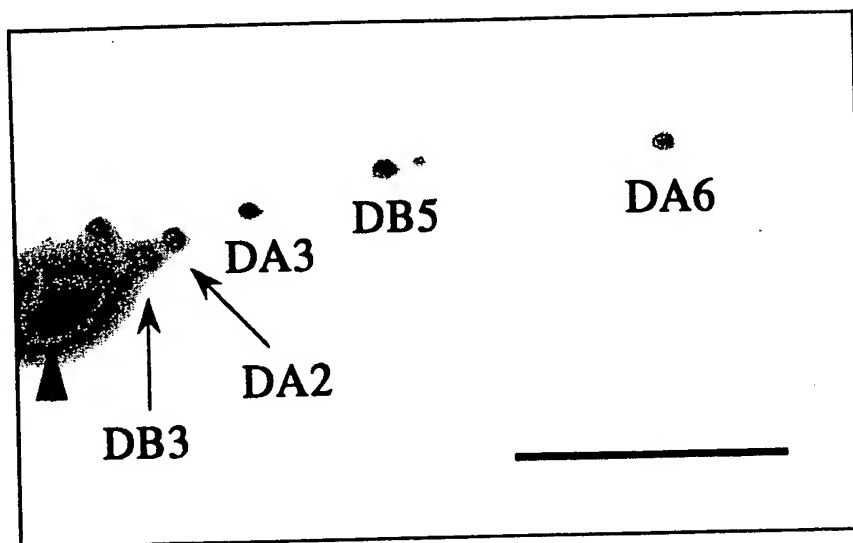


Fig. 6

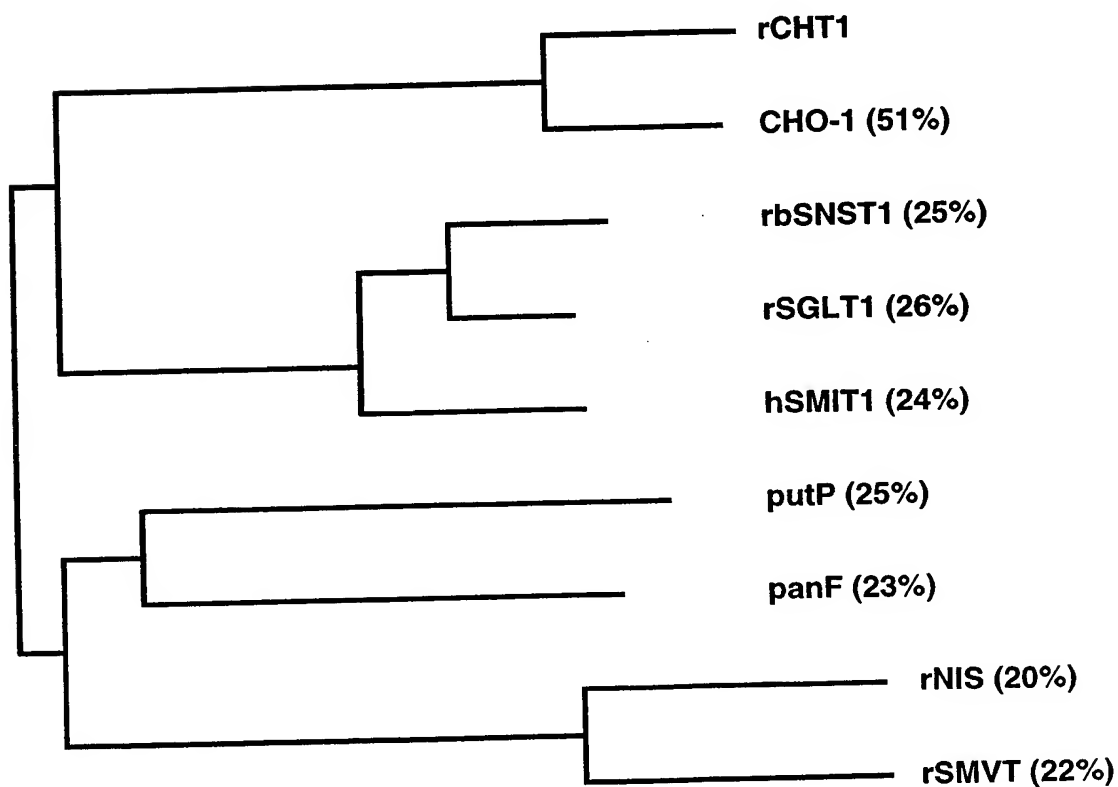


Fig. 7

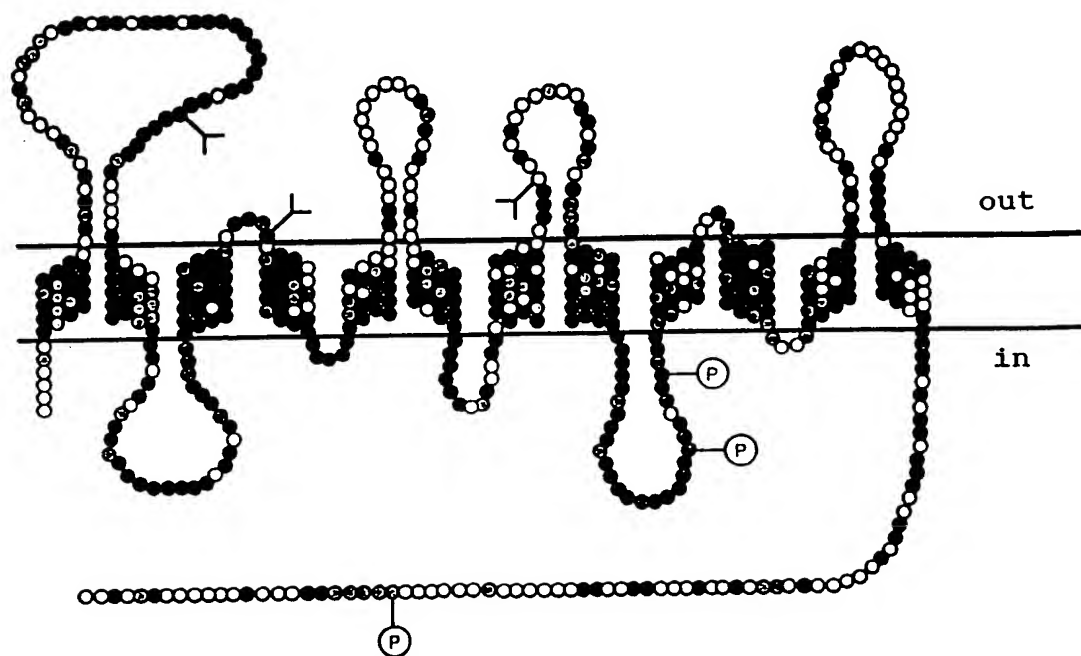


Fig. 8

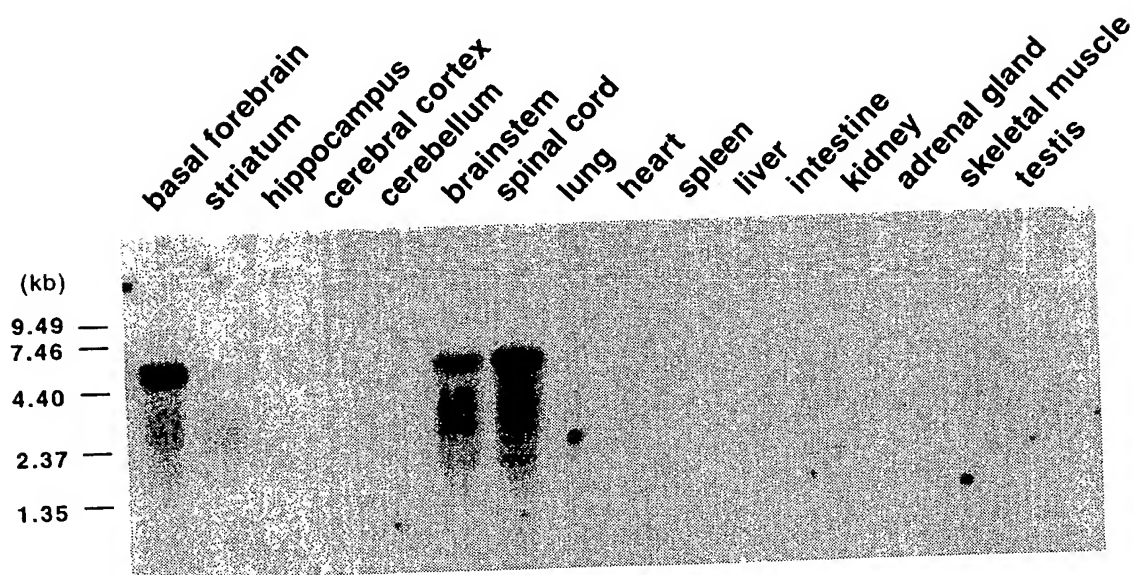


Fig. 9

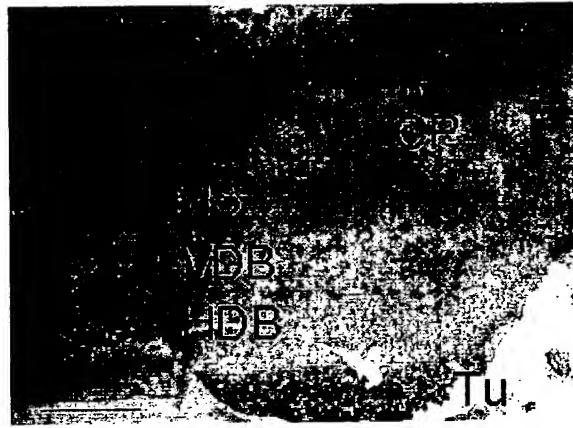


Fig. 10

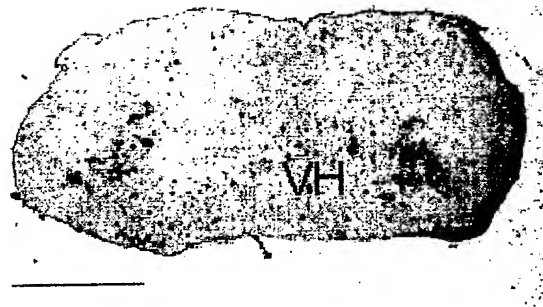


Fig. 11

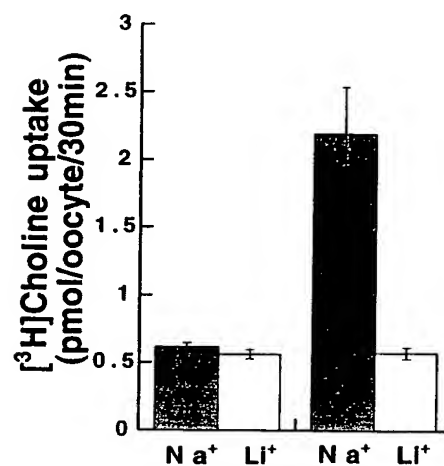


Fig. 12

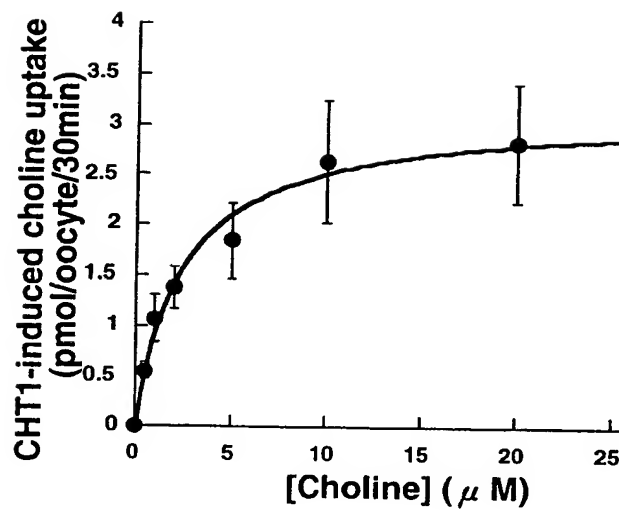


Fig. 13

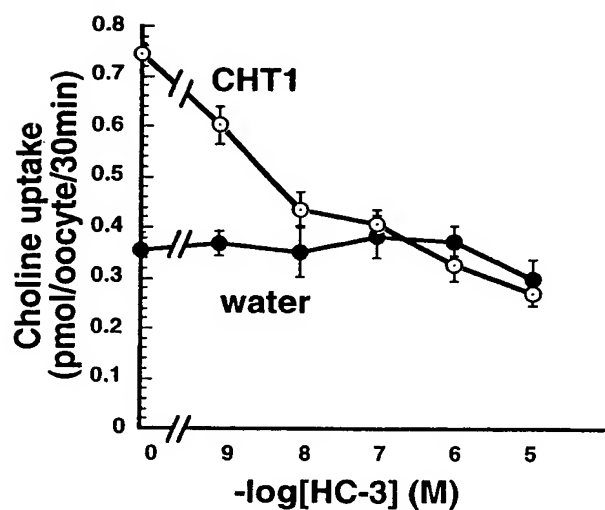


Fig. 14

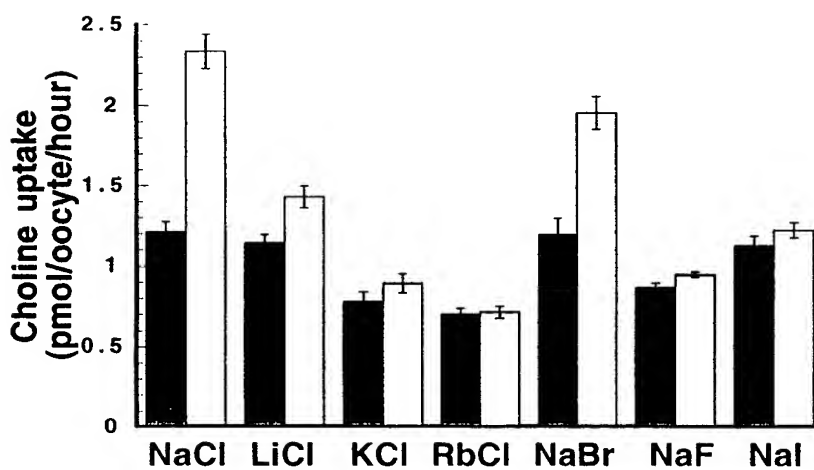




Fig. 15

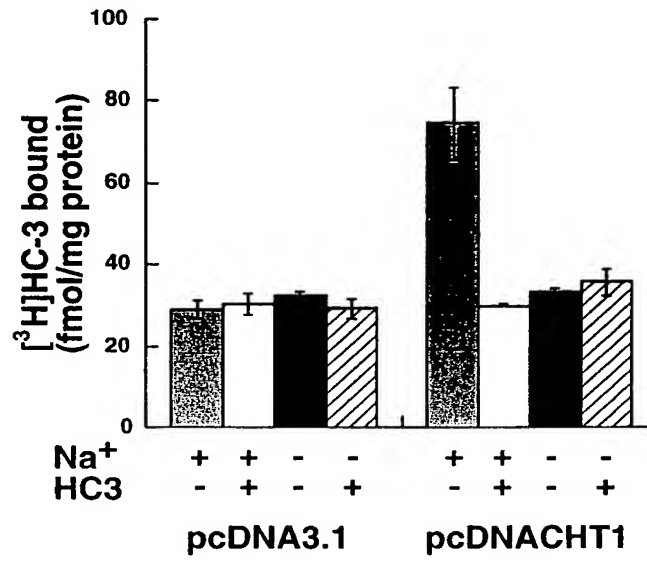


Fig. 16

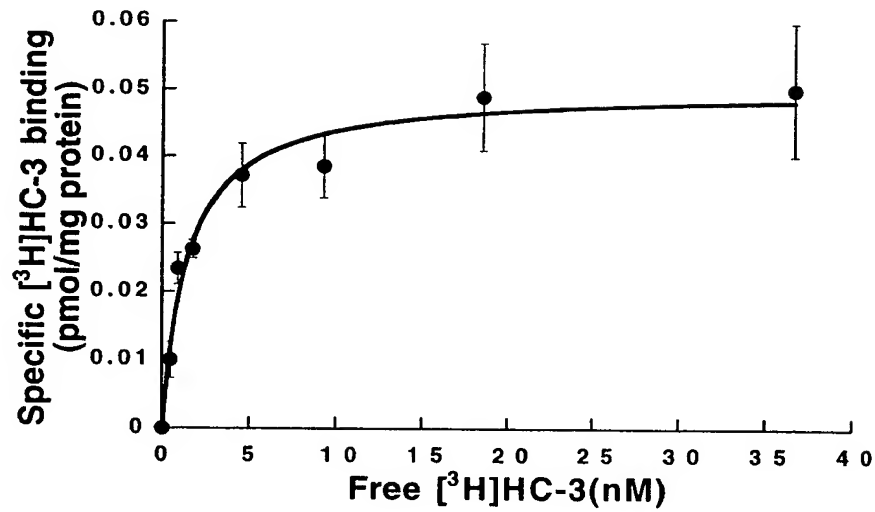


Fig. 17

